#5

FIG. 1A

GCTCCGTGAAGGCCCACCTGCTCTTTCCCAGCTCTGCCGCCTGTGTCTCGGAAGTGTCTCGGTCGAGCA GTAGAAATGCTCAGGGCAAAAGTGCGCTTGATCTGGCGGCTCCAAAAAGCAGCGTGGAGCAGCACTCTT CTGGCCTCGCCCATGAGGCAGTGAGGGTCACAGAGAGTGCATGGAGATCCTGCTGGCAAATA ATGTTAACATTGACCATGCCTCAGCTCGGAACTCCCCTATATGTGGCCTGCACCTACCAGAGGGT CTGCAGCGGCAGTGCTGTGTGTGCTGCTGCAGTTCGGAGCCCAAGGCCCAGTTGGAGGTGCAC CATGCTGCAGCGAGGTCCAGTCATCCACCTGCTAACCGACTATGGAGCTAACCTGAAGC CCTTATTGGAAAATGGTGCACGTCAATGGAGTGACAGTTCACGGAGCCACCCCCTCTTCAATGCTTG AGACTGTGTGAAGAAACTTCTAGAATTAGGAGCCAGTGTCGACCATGGCCAGTGGCTGGACACCCCACTC CTTGTGACAATTAACCGGGTGTCTTCTCTCCACGAGGCATGCCTTGGAGGTCACGTGGCCTTGTGCCAAAG TCATGAAGCTGCAGCTCAGGGGGGGCGCTTACTTAAAACTTTGCACAAGGTGTCAATGTGAAC AAGAAGCGGCTAGGATAGAGAGATCTATGGTGGAATTTTCAGATTGCTGGGCTGATCCCCACT TICTTTAAGCTTTAAAGTTTTTTTTGGCTCTCTCCTAACCCATTTCTATATCGTCAAAGGAAATAGAA TCAGTGAAATTTTACAGTCCTAGGACCCTATACAGAGCATAAGC ATGGCTTTAAAAACATTTTTATTACAATGTTTGCTACGTTTTTT GCGTCCGCGGACGCGTGGGTTATAAC ORF 75-1046 CDNA >33358

tgatttagttccttcccttccccacttttttttttctttttTcTTGAATAATCTGGTATATT ATTGTATAGACACATATAAATTCAGGTGGATAAGATTATAACAAATGTTAGGTATTCCAAGATATGttct GAAGTAAATCCCCCATGAGCTACACTTGAGGAGTGGRAACTCCTGGTTAGTTTAATGTTCTCATTAA AGAGTGAAAAACAATAGGCTGTTTTTTTTTGATTGATTCGGGATCTTGTTGTTGCCTAAAGTCAACATTCTGAAT GCTGTACCTAGAGTTAGAAGCTCAACAGCTTAGACTCCTAGTATCTTTAAATGAGMTCAGTC ACGATTCCTCTATACCAATAGTCTTCCTGGGAAGATACTTGGAATGACACACAGATTGTTGTCT TGTCATCAAGCCACAGAGCTACATCTGCCAGAGCCACTCGA TTGAAAAAAAAAAAAAAAAAA

FIG. 1B

CVKKLLELGASVDHGQWLDTPLHAAARQSNVEVIHLLTDYGANLKRRNAQGKSALDLAAPKSSVEQALLL EAAAQGRLLALKTLIAQGVNVNLVTINRVSSLHEACLGGHVACAKALLENGAHVNGVTVHGATPLFNACC SGSAACVNVLLEFGAKAQLEVHLASPIHEAVKRGHRECMEILLANNVNIDHEVPQLGTPLYVACTYQRVD MEDGPVFYGFKNIFITMFATFFFKLLIKVFLALLTHFYIVKGNRKEAARIAEEIYGGISDCWADRSPLH REGPPALSQLCRLCVRKCLGRACHQAIHKLHLPEPLERFLLYQ. amino acids >33358

FIG. 2A-1

L	-	7	က	4	2	9	7	8	6	10	11 Testes	12 Placenta
	HMVECL	Fetal Brain	Bronchial Epithelium Mix (BEAS-	Mesangial	Fetal Heart	LPS 24 hr Osteoblasts	A2780 WT	Lung	Erythroleuke mia cells	Trachea	(random- primed)	(random- primed)
	HUVECL	Bronchial Epithelium	Cong. Heart Failure	T24 Ct1	Mammary Gland	Burkitt's Lymphoma	A2780 ADR	UCLA-S Lung Carcinoma	Embryonic Keratino- cytes	ME180 IL-1 Cervical Carcinoma	Testes	Mammary Gland (random- primed)
	S/097H	Astrocytes	Cerebellum	T24 Treated	Nat. Killer Cells	Mammary Epithelium	Fetal	p65 Con+/+	SCC25 CDDP- Tongue Squamous Cell Carcinoma	ME 180 Control	RAJI (Burkitt's Lymphoma B cell)	Small Intestine (random-primed)
	U937/ A10P1 0	Prostate Epithelium	Pituitary	Prostate Fibroblast	Cong. Heart Failure (CHFd)	Uterine Smooth Muscle, treated	Esophagus	p65 IL-1+/+	SCC25 WT- Tongue Squamous Cell Carcinoma	MCP-1 Mast cell line	ST486 (Lymphoma B cell)	Fetal Liver (random- primed)

FIG. 2A-2

1	2	Sittor	4	5 Drostato	6 Imbilical	~	∞	Petal	10	11	12 Skeletal
	Primary	Endo-	Bone	Smooth	Smooth	Fetal Liver	A549 con	Hypo-	HPKII	(Acute Pro-	Muscle
	Osteoblast	thelial	Marrow	Muscle	Muscle,			thalamus		myelocytic	(random-
					treated					Leukemia)	primed)
1										Umbilical	
			Cong.		Bronchial			T cells,	Lung	Smooth	Stomach
	Keratino-	Fetal	Heart	Thyroid	Smooth	Fetal Skin	A549 IL-1	CD3	(random-	Muscle,	(random-
	cytes	Kidney	Failure	•	Muscle			treated	primed)	treated	primed)
			(CHFc)							(random-	
										primed)	
6										Uterine	
					WT	Fetal		Tcells,	Heart	Smooth	Spleen
	Melano-	Fetal Liver	Adrenal	LPS 1hr	LNCap+	Adrenal	Fetal	CD3,	(random-	Muscle,	(random-
	cvtes		Gland	Osteoblasts	Casodex	Gland	Testes	IL-4/IL-10	primed)	treated	primed)
								treated		(random-	
										primed)	
-	Coronary				MT		Pulmonary	T cells,	Fetal Brain		Liver
	Smooth	Fetal Lund	Fetal	LPS 6hr	LNCap+	Midterm	Artery	CD3	(random-	Skin/	(random-
	Muscle		Thymus	Osteoblasts	Testos-	Placenta	Smooth	IFNg/TFNa	primed)	Adipose	primed)
	Cells		•		ferone		Muscle	treated			

FIG. 2B-1

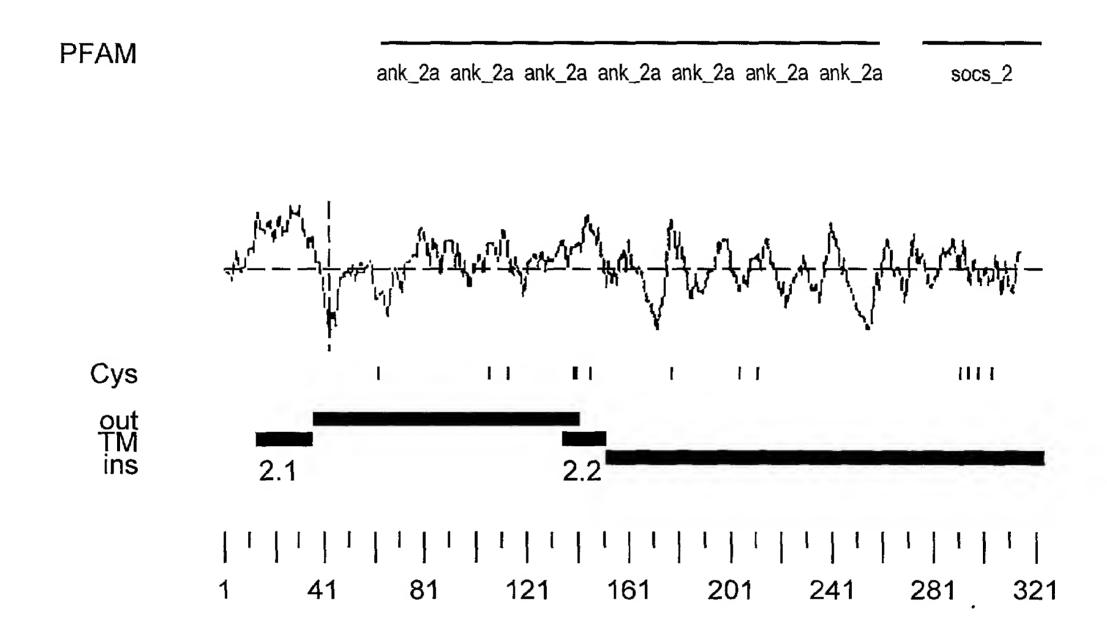
8 9 10 11	spinal HUVEC		(ION3 (umbilical Hypoxia	(ilhbd endo- (umbilical)	thelium)	human Adult Bone	n HUVEC colon Melanoma Marrow	11, control inflamm. G361 cell CD34 +	(jchPa)	W138 20hr HUVEC	334+ serum untreated	Ac) starve Th1 Cells (umbilical	emb. Lung endothelia)	lung	ian squamous IBD Colon	ies cell Th2 Cells (WUM 23)	•
2 9		al erythro-	CHT127	(jthMc)			al CHT1221 skin	n PIT211,	212)	to	r normal BMCD34+	asis breast (jyhMc)	28)	Q.	r ovarian	asis PTH Osteo ascities	
4 5	HMC-1 Fetal	control Dorsal	(mast cell Spinal	line) Cord			Hep-G2 Normal	(insulin- Colon	oma)	Colon to	Normal Liver	Breast Metastasis	Epithelia (CHT128)	Colon to	Normal Liver	Ovarian Metastasis	: :
2 3			Testes Prostate				K563	(RBC line) Uterus				Lung Mammary	Gland			Liver Spleen	
~		A Kidney	(random-	primed)			Brain	(random-	primed)		ら	(random	primed)		D Uterus	(random-	

1 a .

FIG. 2B-2

						• •						
·	_	2	က	4	5	9	7	8	6	10	11	12
						Bun			Prostate			
ш	Thymus			Normal	Colon	adeno-	IBD colon	brain	Tumor	Trigeminal		
	(random-	Heart	Small	Megakaro-	Carcinoma	carcinoma	(MUM6)	subcortical	Xenograft	Ganglia		
	primed)		Intestine	cytes	(NDR109)	(PIT245)	(jbhna)	white	A12			
								matter				
						Bunj			Prostate	_nmpo-		
L	9 week		Retinal	Bone	Colon	sdnamons	Cervical	normal	Tumor	sacaral		
	Fetus	Thymus	Pigmentosa	Marrow	Carcinoma	Sell Sell	Cancer	prostate	Xenograft	Spinal		
		·	Epithelia		(NDR103)	carcinoma	(johCe)	(xoldiz)	K10	Chord		
						PIT299						
									Prostate	-oqmn7		
ග	A549			Ţ-	Colon	d8 dendritic	Spinal cord	ovarian	Cancer to	sacaral		
-	control	Stomach	Retina	induced	Carcinoma	cells	(jlhbc)	epithelium	Liver	Dorsal Root		
-,	(random-			Teell	(NDR82)			tumor	Metastasis	Ganglia		
	primed)								JHH3			
								-T DEANH	Prostate			
I	Salivary	Skeletal	HMC-1	Th-2	Colon	Megakary-	DRG	name	Cancer to	Dorsal Root		
	Gland	Muscle	(mast cell	induced	Carcinoma	ocy	(ION6,7,8)	(umbilical	Liver	Ganglia		
		· · · · · ·	line)	Tcell	(NDR097)	tes(jtheb)	(ilhxe)	end.)	Metastasis	NOI)		
									JHH4	6,7&8)		

FIG. 3



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MEDGPVFYGFKNIFITMFATFFFFKLLIKVFLALLTHFYIVKGNRKEAARIAEEIYGGISDCWADRSPLH
EAAAQGRLLALKTLIAQGVNVNLVTINRVSSLHEACLGGHVACAKALLENGAHVNGVTVHGATPLFNACC
SGSAACVNVLLEFGAKAQLEVHLASPIHEAVKRGHRECMEILLANNVNIDHEVPQLGTPLYVACTYQRVD
CVKKLLELGASVDHGQWLDTPLHAAARQSNVEVIHLLTDYGANLKRRNAQGKSALDLAAPKSSVEQALLL
REGPPALSQLCRLCVRKCLGRACHQAIHKLHLPEPLERFLLYQ.